SEQUENCE LISTING

- <110> Diamandis, Eleftherios P. Kishi, Tadaaki
- <120> Methods for Detecting Ovarian Cancer
- <130> 11757.104USWO
- <140> US 10/510,321
- <141> 2004-10-04
- <150> PCT/CA03/00495
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- <150> US 60/370,559
- <151> 2002-04-04
- <160> 4
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- <210> 1
- <211> 164
- <212> PRT
- <213> Homo sapiens
- <400> 1
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- Leu Leu Gly Gly Ala Trp Ala Gly His Ser Arg Ala Gln Glu Asp Lys
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- Val Leu Gly Gly His Glu Cys Gln Pro His Ser Gln Pro Trp Gln Ala 35 40 45
- Ala Leu Phe Gln Gly Gln Gln Leu Leu Cys Gly Gly Val Leu Val Gly 50 55 60
- Gly Asn Trp Val Leu Thr Ala Ala His Cys Lys Lys Pro Lys Tyr Thr 65 70 75 80
- Val Arg Leu Gly Asp His Ser Leu Gln Asn Lys Asp Gly Pro Glu Gln 85 90 95
- Glu Ile Pro Val Val Gln Ser Ile Pro His Pro Cys Tyr Asn Ser Ser 100 105 110

Asp Val Glu Asp His Asn His Asp Leu Met Leu Leu Gln Leu Arg Asp 115 120 125

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Cys Thr Gln Pro Gly Gln Lys Cys Thr Val Ser Gly Trp Gly Thr Val 145 150 155 160

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Val Leu Gly Gly His Glu Cys Gln Pro His Ser Gln Pro Trp Gln Ala 35 40 45

Ala Leu Phe Gln Gly Gln Gln Leu Leu Cys Gly Gly Val Leu Val Gly 50 55 60

Gly Asn Trp Val Leu Thr Ala Ala His Cys Lys Lys Pro Lys Tyr Thr 65 70 75 80

Val Arg Leu Gly Asp His Ser Leu Gln Asn Lys Asp Gly Pro Glu Gln 85 90 95

Glu Ile Pro Val Val Gln Ser Ile Pro His Pro Cys Tyr Asn Ser Ser 100 105 110

Asp Val Glu Asp His Asn His Asp Leu Met Leu Leu Gln Leu Arg Asp 115 120 125

Gln Ala Ser Leu Gly Ser Lys Val Lys Pro Ile Ser Leu Ala Asp His 130 135 Cys Thr Gln Pro Gly Gln Lys Cys Thr Val Ser Gly Trp Gly Thr Val 145 Thr Ser Pro Arg Glu Asn Phe Pro Asp Thr Leu Asn Cys Ala Glu Val 165 Lys Ile Phe Pro Gln Lys Lys Cys Glu Asp Ala Tyr Pro Gly Gln Ile 185 Thr Asp Gly Met Val Cys Ala Gly Ser Ser Lys Gly Ala Asp Thr Cys 200 Gln Gly Asp Ser Gly Gly Pro Leu Val Cys Asp Gly Ala Leu Gln Gly 210 215 220 Ile Thr Ser Trp Gly Ser Asp Pro Cys Gly Arg Ser Asp Lys Pro Gly 225 230 235 240 Val Tyr Thr Asn Ile Cys Arg Tyr Leu Asp Trp Ile Lys Lys Ile Ile 245 255 Gly Ser Lys Gly 260 <210> 3 <211> 493 <212> DNA <213> Homo sapiens atgggacgcc cccgacctcg tgcggccaag acgtggatgt tcctgctctt qctggggga 60 gcctgggcag gacactccag ggcacaggag gacaaggtgc tggggggtca tgagtgccaa 120 ccccattcgc agccttggca ggcggccttg ttccagggcc agcaactact ctgtggcggt 180 gtccttgtag gtggcaactg ggtccttaca gctgcccact gtaaaaaacc gaaatacaca 240 gtacgcctgg gagaccacag cctacagaat aaagatggcc cagagcaaga aatacctgtg 300 gttcagtcca tcccacaccc ctgctacaac agcagcgatg tggaggacca caaccatgat 360 ctgatgcttc ttcaattgcg tgaccaggca tccctggggt ccaaagtgaa gcccatcagc 420

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